

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:

101581,810

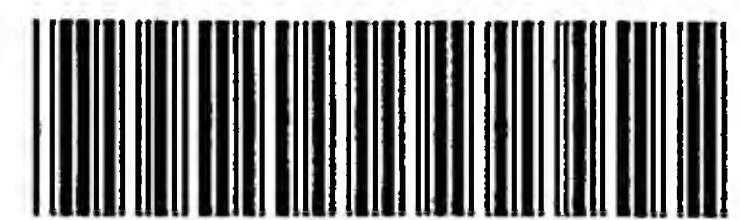
Source:

EFW

Date Processed by STIC:

3/14/07

ENTERED



IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/581,810

DATE: 03/14/2007

TIME: 14:58:16

Input Set : N:\efs\03_14_07\10581810_efs\HERR13001APCSEQLIST.TXT
 Output Set: N:\CRF4\03142007\J581810.raw

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4 <110> APPLICANT: Ortega Mora, Luis MIguel
5     Fernandez Garcia, Aurora
8 <120> TITLE OF INVENTION: Use of gene NcSAG4 for the diagnosis and
9     prevention of neosporosis and as a marker for analysis of
10    the pathogenesis
12 <130> FILE REFERENCE: HERR13.001APC
14 <140> CURRENT APPLICATION NUMBER: US 10/581,810
15 <141> CURRENT FILING DATE: 2006-06-02
17 <150> PRIOR APPLICATION NUMBER: PCT/ES2004/000529
18 <151> PRIOR FILING DATE: 2004-11-26
20 <150> PRIOR APPLICATION NUMBER: ES P200302869
21 <151> PRIOR FILING DATE: 2003-12-04
23 <160> NUMBER OF SEQ ID NOS: 15
25 <170> SOFTWARE: FastSEQ for Windows Version 4.0
27 <210> SEQ ID NO: 1
28 <211> LENGTH: 23
29 <212> TYPE: DNA
30 <213> ORGANISM: Neospora caninum
32 <220> FEATURE:
33 <221> NAME/KEY: modified_base
34 <222> LOCATION: 6
35 <223> OTHER INFORMATION: I
37 <400> SEQUENCE: 1
W--> 38 tggacntayg aytyaaraaa rgc                                23
40 <210> SEQ ID NO: 2
41 <211> LENGTH: 23
42 <212> TYPE: DNA
43 <213> ORGANISM: Neospora caninum
45 <220> FEATURE:
46 <221> NAME/KEY: modified_base
47 <222> LOCATION: 12, 15, 18, 21
48 <223> OTHER INFORMATION: I
50 <400> SEQUENCE: 2
W--> 51 aaraargara tnatnacncc ngg                                23
53 <210> SEQ ID NO: 3
54 <211> LENGTH: 23
55 <212> TYPE: DNA
56 <213> ORGANISM: Neospora caninum
58 <220> FEATURE:
59 <221> NAME/KEY: modified_base
60 <222> LOCATION: 3
61 <223> OTHER INFORMATION: I
63 <400> SEQUENCE: 3

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W--> 64 acnggytcrt cyttrcarty rtc	23
66 <210> SEQ ID NO: 4	
67 <211> LENGTH: 24	
68 <212> TYPE: DNA	
69 <213> ORGANISM: Neospora caninum	
71 <220> FEATURE:	
72 <221> NAME/KEY: modified_base	
73 <222> LOCATION: 22	
74 <223> OTHER INFORMATION: I	
76 <220> FEATURE:	
77 <221> NAME/KEY: misc_feature	
78 <222> LOCATION: 7	
79 <223> OTHER INFORMATION: n = A, T, C or G	
81 <400> SEQUENCE: 4	
W--> 82 rtcytnacy ttraarcara angg	24
84 <210> SEQ ID NO: 5	
85 <211> LENGTH: 20	
86 <212> TYPE: DNA	
87 <213> ORGANISM: Neospora caninum	
89 <400> SEQUENCE: 5	
90 ccgacgaagc cctgagaact	20
92 <210> SEQ ID NO: 6	
93 <211> LENGTH: 19	
94 <212> TYPE: DNA	
95 <213> ORGANISM: Neospora caninum	
97 <400> SEQUENCE: 6	
98 tgtcgctgt tgggttgt	19
100 <210> SEQ ID NO: 7	
101 <211> LENGTH: 24	
102 <212> TYPE: DNA	
103 <213> ORGANISM: Neospora caninum	
105 <400> SEQUENCE: 7	
106 gaaacaagaa aagagactat ctca	24
108 <210> SEQ ID NO: 8	
109 <211> LENGTH: 20	
110 <212> TYPE: DNA	
111 <213> ORGANISM: Neospora caninum	
113 <400> SEQUENCE: 8	
114 ccaggtgaga gtgtttcgat	20
116 <210> SEQ ID NO: 9	
117 <211> LENGTH: 601	
118 <212> TYPE: DNA	
119 <213> ORGANISM: Neospora caninum	
121 <220> FEATURE:	
122 <221> NAME/KEY: CDS	
123 <222> LOCATION: (48)...(568)	
125 <400> SEQUENCE: 9	
126 ggcaaacacgt cgcaacgtac tctcatctt ttctgtgggtt tgcagcc atg gag aaa	56
	Met Glu Lys
127	

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128		1		
130	agc gcc ttc ttt ccc agg gtg gtg ctc tgt ttc gtc gta gtt ttg tcc		104	
131	Ser Ala Phe Phe Pro Arg Val Val Leu Cys Phe Val Val Val Leu Ser			
132	5	10	15	
134	gcg tgc tcg gcg tgg cga gtg gaa ggg aag aac tgg tcg tac gat ttc		152	
135	Ala Cys Ser Ala Trp Arg Val Glu Gly Lys Asn Trp Ser Tyr Asp Phe			
136	20	25	30	35
138	aag aag ccg ctg gac agc gat gaa aca aga aaa gag act atc tca cca		200	
139	Lys Lys Pro Leu Asp Ser Asp Glu Thr Arg Lys Glu Thr Ile Ser Pro			
140	40	45	50	
142	ggt gag agt gtt tcg ata caa aat tct ggg agc att acg ctg gcg tac		248	
143	Gly Glu Ser Val Ser Ile Gln Asn Ser Gly Ser Ile Thr Leu Ala Tyr			
144	55	60	65	
146	aac cca aca ggc gac aca caa gtt ctc agg gct tcg tcg gga gac agc		296	
147	Asn Pro Thr Gly Asp Thr Gln Val Leu Arg Ala Ser Ser Gly Asp Ser			
148	70	75	80	
150	tgc agg gat gag cca atc gaa ctt gcg act tta ttc cca gca gcc acg		344	
151	Cys Arg Asp Glu Pro Ile Glu Leu Ala Thr Leu Phe Pro Ala Ala Thr			
152	85	90	95	
154	ccg gcg ccc acg tgg atg caa act ggt agc acg aga acc tta gcg ttt		392	
155	Pro Ala Pro Thr Trp Met Gln Thr Gly Ser Thr Arg Thr Leu Ala Phe			
156	100	105	110	115
158	cct acc aac gca gta ccc gcg aag cag acc acg ccg ttc tgt ttt aaa		440	
159	Pro Thr Asn Ala Val Pro Ala Lys Gln Thr Thr Pro Phe Cys Phe Lys			
160	120	125	130	
162	gtc acg gat acg cag aag aac aaa act ctg aca gcg ata atc aag gtc		488	
163	Val Thr Asp Thr Gln Lys Asn Lys Thr Leu Thr Ala Ile Ile Lys Val			
164	135	140	145	
166	gcc ggt gcc caa ggc ttg tct gct gct ctg ggg gtc tcc att gga ata		536	
167	Ala Gly Ala Gln Gly Leu Ser Ala Ala Leu Gly Val Ser Ile Gly Ile			
168	150	155	160	
170	cca gct ctt gct ttt gca ctg agt tcg ata ta agggcatgca aacgaataaa		588	
171	Pro Ala Leu Ala Phe Ala Leu Ser Ser Ile			
172	165	170		
174	tgagggcgact gat		601	
176	<210> SEQ ID NO: 10			
177	<211> LENGTH: 173			
178	<212> TYPE: PRT			
179	<213> ORGANISM: Neospora caninum			
181	<400> SEQUENCE: 10			
182	Met Glu Lys Ser Ala Phe Phe Pro Arg Val Val Leu Cys Phe Val Val			
183	1	5	10	15
184	Val Leu Ser Ala Cys Ser Ala Trp Arg Val Glu Gly Lys Asn Trp Ser			
185	20	25	30	
186	Tyr Asp Phe Lys Lys Pro Leu Asp Ser Asp Glu Thr Arg Lys Glu Thr			
187	35	40	45	
188	Ile Ser Pro Gly Glu Ser Val Ser Ile Gln Asn Ser Gly Ser Ile Thr			
189	50	55	60	
190	Leu Ala Tyr Asn Pro Thr Gly Asp Thr Gln Val Leu Arg Ala Ser Ser			

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191	65	70	75	80
192	Gly Asp Ser Cys Arg Asp Glu Pro Ile Glu Leu Ala Thr Leu Phe Pro			
193	85	90	95	
194	Ala Ala Thr Pro Ala Pro Thr Trp Met Gln Thr Gly Ser Thr Arg Thr			
195	100	105	110	
196	Leu Ala Phe Pro Thr Asn Ala Val Pro Ala Lys Gln Thr Thr Pro Phe			
197	115	120	125	
198	Cys Phe Lys Val Thr Asp Thr Gln Lys Asn Lys Thr Leu Thr Ala Ile			
199	130	135	140	
200	Ile Lys Val Ala Gly Ala Gln Gly Leu Ser Ala Ala Leu Gly Val Ser			
201	145	150	155	160
202	Ile Gly Ile Pro Ala Leu Ala Phe Ala Leu Ser Ser Ile			
203	165	170		
206	<210> SEQ ID NO: 11			
207	<211> LENGTH: 26			
208	<212> TYPE: DNA			
209	<213> ORGANISM: Neospora caninum			
211	<400> SEQUENCE: 11			
212	gccaggatcc atggagaaaa gcgcct			26
214	<210> SEQ ID NO: 12			
215	<211> LENGTH: 31			
216	<212> TYPE: DNA			
217	<213> ORGANISM: Neospora caninum			
219	<400> SEQUENCE: 12			
220	ttcgaattcc ttatatcgaa ctcagtgcaa a			31
222	<210> SEQ ID NO: 13			
223	<211> LENGTH: 28			
224	<212> TYPE: DNA			
225	<213> ORGANISM: Neospora caninum			
227	<400> SEQUENCE: 13			
228	ttatggatcc ggaagaactg gtcgtacg			28
230	<210> SEQ ID NO: 14			
231	<211> LENGTH: 30			
232	<212> TYPE: DNA			
233	<213> ORGANISM: Neospora caninum			
235	<400> SEQUENCE: 14			
236	tttgaattcc ttaggcgacc ttgattatcg			30
238	<210> SEQ ID NO: 15			
239	<211> LENGTH: 155			
240	<212> TYPE: PRT			
241	<213> ORGANISM: Neospora caninum			
243	<400> SEQUENCE: 15			
244	Met Arg Gly Ser His His His His His His Gly Met Ala Ser Met Thr			
245	1	5	10	15
246	Gly Gly Gln Gln Met Gly Arg Asp Leu Tyr Asp Asp Asp Asp Lys Asp			
247	20	25	30	
248	Arg Trp Ile Arg Lys Asn Trp Ser Tyr Asp Phe Lys Lys Pro Leu Asp			
249	35	40	45	
250	Ser Asp Glu Thr Arg Lys Glu Thr Ile Ser Pro Gly Glu Ser Val Ser			

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251 50 55 60
252 Ile Gln Asn Ser Gly Ser Ile Thr Leu Ala Tyr Asn Pro Thr Gly Asp
253 65 70 75 80
254 Thr Gln Val Leu Arg Ala Ser Ser Gly Asp Ser Cys Arg Asp Glu Pro
255 85 90 95
256 Ile Glu Leu Ala Thr Leu Phe Pro Ala Ala Thr Pro Ala Pro Thr Trp
257 100 105 110
258 Met Gln Thr Gly Ser Thr Arg Thr Leu Ala Pro Thr Asn Ala Val Pro
259 115 120 125
260 Ala Lys Gln Thr Thr Pro Phe Cys Phe Lys Val Thr Asp Thr Gln Lys
261 130 135 140
262 Asn Lys Thr Leu Thr Ala Ile Ile Lys Val Ala
263 145 150 155

RAW SEQUENCE LISTING ERROR SUMMARY
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Input Set : N:\efs\03_14_07\10581810_efs\HERR13001APCSEQLIST.TXT
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 6
Seq#:2; N Pos. 12,15,18,21
Seq#:3; N Pos. 3
Seq#:4; N Pos. 7,22

VERIFICATION SUMMARY
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Input Set : N:\efs\03_14_07\10581810_efs\HERR13001APCSEQLIST.TXT
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L:38 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:64 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:82 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0